

Figure 1

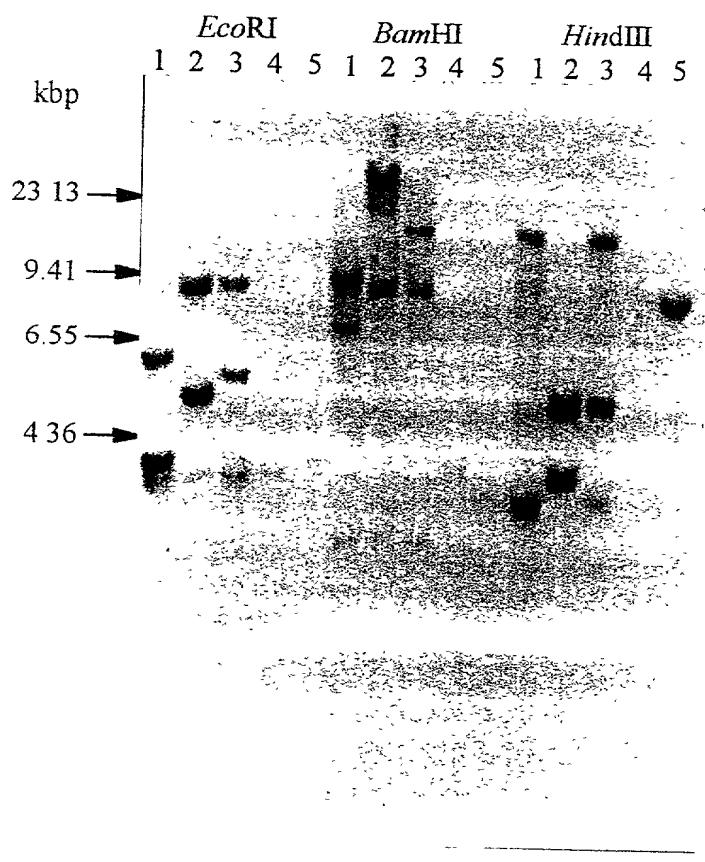


Figure 2

1 CAGCATTCCA AGAGGAAAAA AAACATGATC AAGAAGTAAT TACTACAAA
51 GAGGAAGCTG TAGTAGTAAC TGCACCACCA CCATCAGAAA CAGCAGAGCC
101 AGCTGCAGCT GTTGTGCGG AGGAAGAGAC AACAAAGGAG CAAGAAGAGC
151 CGCCAGCAGT ATCGGCCGAG GAACCTGTGG CCCCAGCTGA AGTAGAGACA
201 AAGGTGGAAG TTACAGAAGA ACCACCAAAA GTTGAGGAGA AACCAGCAGA
251 AGTAGAGGAG GCTCCAAAGG AAACAGTAGA AACAGAACCA GCTGTTGAGA
301 AGACCATCAA GGAGGAAACT GTAGAGGACT CTGTCGTGGC ACCTGCTCCC
351 GAACCGGAAG CCGAAGTCCC AAAAGAGAAG GTAATTGCTA CTACTGAAAC
401 TACTGAGGAA GAAGAAAAAG TGGCAGTTGA AGAAGTTGAA GTGAAAGTTG
451 AAACAGAGGA GGGAGAAGTT ACTGAGGAGA AGACTGAGTA AAATAAGTTG
501 TACAACTATT TTATGCACGC CTTATTTCT CAAATTGGAAG TTTATAATGT
551 AGTGGGCTTT TGGTAATATT TGGGGGTTA ATAAGTGGTT TAAGTGGTT
601 AAGGCTTTT TGGAATTTAG ATATTTGGGT AAAGGCCTAC TTGAACAAAA
651 CATAGAAATT TGGCACACAT GGGTAAAAGT CAAACTTGT TGAGGATGTT
701 TTCTTGTG TGAAATGTGT GTGCCAAGTA GTAGAATGTG GTGGTTGAA
751 TGTAAGTTCT CAAGTAGGGT TTATGAGTCC TAGTATTATG CTTGATTGTA
801 TGTTGATATG AAAATGGGGG TATGTTGGCT TTGAATAAAA GTTTTTAATT
851 TTATAAAAAA AAAAAAAAAA AAAAAAAAAA AA

Figure 3

1 AFQEEKKHDQ EVITTKEEAV VVTAPPSET AEPAAAVVAE EETTKEQEEP
51 PAVSAEEPVA PAEVETKVEV TEEPPKVEEK PAEVEEAPKE TVETERAVEK
101 TIKEETVEDS VVAPAPEPEA EVPKEKVIAT TETTEEEEKV AVEEVEVKVE
151 TEEGEVTEEK TE

Figure 4

1 AAACAACAAACTTTTCAATCTTCTTCATCACCATGTCGAGCTGCGGAA 60
T T N F F I N L L S L I I T M S S C G N

61 ACTGCGACTGTGCCGACAAGACCAACTGCCCAAAGAAGGGAAACAGCTACGGCTTGACA 120
C D C A D K T N C P K K G N S Y G F D I

121 TCATTGAGACCCAGAAGAGCTACGATGACGTGCGGTGATGGATGTTCAGGCAGCTGAGA 180
I E T Q K S Y D D V V V M D V Q A A E N

181 ATGATGGCAAGTGCAGTGCAGGGCCCGAGCTGCAGTTGTGATGGCTGCAGCTGTGGTCATT 240
D G K C K C G P S C S C V G C S C G H *

241 AAGTTAACACAAACATTATCATGTTATAGTGAATAATGATGTGTGATGAATATAGGTG 300
301 AAAATCTGTGGTGTGATAAAAACCGTTGGTGAATAAAATAGGTGTATATTCGTGTGCAC 360
361 CTTCTACGAGTACTTGTGCTTGGGTGAAAGAAATATGCACCTAACGTGTCAGTTGTT 420
421 TCCGTGTTTCGCCGTGTCCTTGTAAATGGTCATGTTGTGTTCTGTGGTTAAATT 480
481 AAATGAACTAGTAATGTTATGTAAAAAAAAAAAAAAA 519

Figure 5

1 GGAGGAGATCACCAACAGTCGTCTAATGAGACACGGCGATGGATAGAC 60
 R R S P V P P T R R R N E T R R S D R Q

 61 AACTTCGAGCCACTGTGGGTGAAGACGGCGCGAACGATGGGACCCACCCCTGGTCATG 120
 L R A T V G E D G G E R W D P P L V D E

 121 AAGGCAAGCTCCGTACCTTCCGGACAGGTCTGAAGCTCCGAACCAATTGATTTCCGA 180
 G K L R T F R T G L K L R T N F D F P I

 181 TCCATCGTGTCTTGATCACCTTCCGTGCGTACAGACAGCATCGGAAGTCATCT 240
 H R V F V S P F L R C V Q T A S E V I S

 241 CCGCTCTCTGCAGCGTCGACGATATTCCGCCACCACTAATAGAGGCATCAAGTACAAA 300
 A L C A V D D I P A T T N R G D Q V Q I

 301 TCGATCCATCCAAGATCAAGGTCTCTATTGAGTATGGATTATGTGAAATGTTAACATGC 360
 D P S K I K V S I E Y G L C E M L N M Q

 361 AAGCCATAAGACTTGGTATGGATTCAGCAATGGGAATTGGGTTTCGATAAAATCACACC 420
 A I R L G M D F S N G N W G F D K S H L

 421 TTGAATCAACATTCCCAGTTGGGACGGTGGATCATAGTGTGAAACCACTCTATAAGAGA 480
 E S T F P V G T V D H S V E P L Y K E M

 481 TGCCAAAATGGGAAGAGACAGTCAATGGCGCAAGGGCCAGATATGAAGAGGTTATTCAAGG 540
 P K W E E T V N G A R A R Y E E V I Q A

 541 CCCTAGCAGATAAAATACCCACGGAGAACTTGGCTTACACATGGGAAGGAGTG 600
 L A D K Y P T E N L L L V T H G E G V G

 601 GCGTTGCAGTTCTGCCTTCATGAAGGATGTTACAGTGTACGAAGCCGATTATTGTGCCT 660
 V A V S A F M K D V T V Y E A D Y C A Y

 661 ATACACACGCAAGAAGATCCATTGTCTTGGGCAAAACCCAGTCATTTACTGCTGAAACT 720
 T H A R R S I V L G K N Q S F T A E N F

 721 TTGAAGTATTACCAAAACAAGGCCAAACTGGTGTCAAGTTACGTCTTGAACAGCATTGAT 780
 E V L P K Q G Q T G V S Y V L E Q H *

 781 GGAACGTATGACCTAATTGTGGCAGCCGATGATTACAGAAACAATTCCACACCTTTT 840
 841 TCTTTTTTCGGGCATTGCCTACATTTATAATTAAATTAGGCATTCTCATAGCTAAGGCT 900
 901 CATTGGATTTCACATCCCTACTTGTAAAGGGAGACTTGATTTGCTTGCCTCAAACAGAA 960
 961 CATATGTTGCTGTGTCATCAGCTTTTTAACTGGGATTTCTATTTTACAGTGTGTAA 1020
 1021 AAAAAAAAAAAAAAAA 1046

Figure 6

1 GTTGATGGCAGATGTGACCAACTCAGGAAAAATGCCAGGGTTGCAATTGATTCTTAC 60
 V D G R C D Q L R K N A R V V A I D S Y

 61 GAAGATGTTCTTGAACGATGAGAACGCATTGAAAAAGGCAGTGGCTAGTCAGCCTGTG 120
 E D V P L N D E N A L K K A V A S Q P V

 121 CGCGTCGCCATTGAAGGAGGTGGCAGGGATTCACACTCTATCAATCAGGCCTTTACT 180
 R V A I E G G G R D F Q L Y Q S G V F T

 181 GGATCATGTGGGACGGCCCTAGACCATGGTGTGGCTGCTGGGTATGGCACAGAAAAT 240
 G S C G T A L D H G V A A V G Y G T E N

 241 GGTGTGGATTACTGGATTGTAAGGAACCTCATGGGTGCAAGCTGGGAGAGAGCGGCTAC 300
 G V D Y W I V R N S W G A S W G E S G Y

 301 ATCAGGATGGAACGTAATCTGGCAGGCACAGCTACGGCAAATGTGGTATTGCAATGGAA 360
 I R M E R N L A G T A T G K C G I A M E

 361 GCCTCTTACCCATTAAAGAAAGGCCAAATCCCCAAACCCAGGACCATCTCCCTCCATCT 420
 A S Y P I K K G Q N P P N P G P S P P S

 421 CCAATAAAAGACCTCCAACAGTTTGACAATTACTATACCTGGCTGAAAGCACCCTT 480
 P I K T S N S F V T I T I P W L K A P L

 481 GCTGCTGTCTATTGAGTTGGCAGGTATTGCTTCGAGTGGGATGTTGCCACTCGAGG 540
 A A V Y L S L A G I A S S G D V A H S R

 541 CTGCCACTTGCTGTGATGACCATTACAGTTGCTGCCACATGAGTATCCCCTGCAACC 600
 L P L A V M T I T V A A H M S I P S A T

 601 TTAATGCAGGGACGTGTATGATGAGAAGGACAACCCATTGAGTGTGAAGGCATTGAAGCG 660
 L M Q G R V *

 661 TACTCCGCTAACCTCATTGGCCTTGGGAACCGTGGCAAGAGCAGCAGTGCTTAAGA 720
 721 ACATTGTGTCATCTATACAGTGAAGTAAACGAGGATGAAAAGTTGTATCAGGCAGGGC 780
 781 TTGATGATCTCCTCGGTTTATAGTACCGCATAACCTCATTCTCCATTAGGTATATAC 840
 841 ATATGGACGGTTTATCAAAGTTATTCAAGATGCTATTATGTATATCATTCTCAGTC 900
 901 CCTGTATTCATTAAACGAGAACATAAACAGATCGTTATCAGCTACCAATTCCACTGT 960
 961 AACACGTTATCAATTATTTACTGGCCTCGCTGAAAAAAAAAAAAAAA 1017

Figure 7

1 CGGTTCAATCGCTGGATCAATCGAGCATATGGCGATGTATCCGGTTGATACGCTTAAAC 60
 G S I A G S I E H M A M Y P V D T L K T

 61 TCGCATAACAGGCTATTGGGTCAATGTCGGCTCAATCCGCCGGTCTCGACAAGCCCTGG 120
 R I Q A I G S C S A Q S A G L R Q A L G

 121 GTCGATACTGAAAGTTGAAGGTCCCGCCGGACTTACCGTGGCATTGGTCAATGGGTCT 180
 S I L K V E G P A G L Y R G I G A M G L

 181 CGGTGCAGGACCAGCTCACCGCAGTGATTTCTCGTTACGAGATGTGTAAGGAGACTT 240
 G A G P A H A V Y F S V Y E M C K E T F

 241 TTCTCATGGTATCCGAGCAATTCCGGTGCACGCCGTTGGGGGTGTTCGCGACGGT 300
 S H G D P S N S G A H A V S G V F A T V

 301 GGCAAGCGACCGCGGTGATTACGCCGATGGATGTGGTAAACAGAGGTTGCAGTCAGAG 360
 A S D A V I T P M D V V K Q R L Q L Q S

 361 CAGTCCTGACAACGGGTGTTGTTGATTGCGTGAGGAGGGTGGTAGAAGAAGGGATTGG 420
 S P Y K G V V D C V R R V L V E E G I G

 421 CGCATTTTACGCATCTTATCGAACAACTGTGGTCAATGCCCGTTACGCCGTTCA 480
 A F Y A S Y R T T V V M N A P F T A V H

 481 CTTGCCACATATGAAGCCACGAAGAAAGGGTTGGAGGTGTCGCCGGAGACTGCAA 540
 F A T Y E A T K K G L L E V S P E T A N

 541 CGATGAGAATTGTTAGTCATGCTACTGCTGGTGTGCTGGAGCTTGGCTGCAGT 600
 D E N L L V H A T A G A A A G A L A A V

 601 AGTAACCACTCCACTAGATGTTGCAAAACTCAGTTGCAGTGCAAGGTGTTGGGATG 660
 V T T P L D V V K T Q L Q C Q G V C G C

 661 CGACAGATTTCTAGCAGTCGATTCAGGATGTTAGGAAGCATAGTGAAGAAAAATGG 720
 D R F S S S S I Q D V I G S I V K K N G

 721 ATATGTCGGGTTAATGAGGGGGTGGATTCCCAGAACATGCTATTTCATGCTCCTGCTGCAGC 780
 Y V G L M R G W I P R M L F H A P A A A

 781 AATCTGCTGGCTACTTATGAAGCCTCCAAAACATTCTTCAAAACTCAATGAGAGCAA 840
 I C W S T Y E A S K T F F Q K L N E S N

 841 TAGCAACAGCTCAGTTACCTAACGATTTCATATGTTTGTGCTACTAGGCTTATCCA 900
 S N S S V T *

 901 AAATCATGTCGATTGGTTCACTTCACCACAGTGGCATGAACTCAAAGCATCGAAT 960
 961 TTTACATGTATATTATGCAATCTAGATGCTTCTGATATTTATTTTATTTTCTTTC 1020
 1021 CAACTTTGTAATTAGAATTAGCTACTATGGTTATGGCATGGAGTGTATAATTGCTA 1080
 1081 ATATCATCGTATAAGCAATGCTATTGAGAAATTGGGTGTAAGGTTAGAGTAATGTTAT 1140
 1141 TTGCACAATCCACTACATAGACCGCGGGACTCAAAAAAAAAAAAAAAA 1195

Figure 8

1 GAGCTTATAT TCGCTTTCAAA AGCTTCATAA TTACCTTCAAA TGTAACTCTC AACCTTCTTA AGCTTGTAT CTTATTAACG AACCCACAA CACATTATT
 101 ATGAACTCT TGGACTTTT GCTCTGTAC CAATAATCGC ACCAACAAA AAATTCCTTT TGTTTATAT TCGTTTTTA TTTTTAAC GTTGTGAT
 201 TCAAACATCA TATAGTGAG GGGGANTATT ATTCGGACTC CTCCAAAAC TTATGACATT GTGATTACAC ATTGATGTA CAGAAGTTT TGATGAAGTG
 301 CCAATTCAA TCYTTCCTA ATTGCTTCAT AAAGGGTGT TTGTTAATTA ATAGAAGT AAGGAAATT AGCAAGAAGT GCATTATGG GACTGGTATA
 401 TATGAGGG ATCTGCGCTG GUAAGGAGG AAAGGGTGC CTGAGTCAGG TGTGTCCTAAT CGTGAAATAT TCTTCMAMG AGAGCCACC ATCTCATAG
 501 'YGGATTTAG AAAGTGGTTT CCACAAAAAA ATATGACACA ACCCATCCAT GAAACAAATA AACATGACA GGTCTATCATT TCTTCTCTC
 601 AAGKATAAA TACCTATTAAG TGTCTTAAC ACCGGCCTAA CTTGTCATT CTTGTCATT' GGTGACTTT TATGCCCCAA TTGTTGCTTG AAGGAATAA
 701 AAAGGAAATG CTYYTCCTTG AACCCATATG GAAAGGATT CATGAGGAA GATAGAGGA GATAGAGGG AGGGGGAGG ATYGGGGAG AGAATTGATA CGGAGTCCTP
 801 'TTAAATGGTA TATGTAATTC ACTCAGAAAC ACCTGATACCA TATATGCATC AATGTCAATG TCACAGAAA CGTARCTCAC GAACCATTT CGTAACATGC
 901 ATGCAACCAT CATACTTAT AACATAGTGT TACGACAATA AAAGACCTT AGTGTAAAGA GCATTAGTC GTGACAGAA CAAAACGTG GATGCCAAC
 1001 CTAACGAGG GTATATCTT TATTCATATA TCTACTTGT ATATGACCTA AACCTTGTGT CACCCAAAT GTTCGTAGC ATGCTTATTT GTTIGACTG
 1101 'TGPGGGATGA GAAATGATAT GAGCTGTGCC ATTAATTTA GCGGGATGTG ATTTGGGTAT ATATGACCA ATATAAGATA TATAAAACTT GAACTAAACA
 1201 ATTTTCCTAAC AGTTTAAACT ATAAAGTAAAT CTCCTTCAG ATGTTAAACT ATTTGGGTAG ATATCCGGTG ATTAACCCCA ATTAATTTAA ATCTCCAGCA
 1301 AAATCTGTA TCTCTTCTC TCGAGGGA ATTCCTTCCT TCCANACACC TTAACTAAAG TAGTAAAGT TAGTAAAGT MATTGAAA TGATAACACA
 1401 AGAGTGATA AGGTGATGG TCACTACTT ACCCRACTGC ACAAACACA CAGGCACACA TCCAAAGTA GTAGTATGAT TACACATTT 'GAAAANATG
 1501 ACCUCATTA TTAGGCCAC CTCCTGTAA AAAAGATA CAAACAAATT ACCCTATCA TTATATAA ATTTGAGCA TAACTCTTC TCCAATCCAC
 1601 ACCATTTT TTAATTTT GCAAAACATG CTAAAGCTT CTGTATCA GTCAGAATG GTGTCAATG CCCAAGATC TCACTGCCC CTCCTCTCT
 1701 CTCCTCTCT CTCCTCTCT CCTCTCTCTC TCTCTCTC TCTCTCTC ATCAACTGAA GGCTTTAGG ACCCTTATAT AAACCTCTT CAATGATCA TCTCTG

Figure 9

1 GATCTTATATTGAGGATGCAAAGTTCAAATTACCTGATACTGAACAAATCA 60
 61 AGCTTTGATCATATAATCGAAACCAACACACAATAATTATGAATTCTTGACTCTT 120
 121 GTCTCTGTACCAAAATACGCACACCACAAAAAATTCTTTGTATTATTCGTTTTA 180
 181 TTTTTTAACGTTGGTATTCAAACATCATATAAGTAAGGGGAATATTATTCGGACTC 240
 241 CTCCAAAAACTATGACATTGTGATTACACATTGAATGACAGAAGTTTGATGAAGTG 300
 301 CCAATATCAATCTTCTTAATTGCTCATAAAGGGTGTGTTGTAATTAAAAGAAAGAT 360
 361 AAGGAAATTAGCAAGAAGTGCATTATTGGGACTGGTATATATGACAAGGATCTGACGTG 420
 421 GCAAAGAAAGAAAGTGGGCTCTGAGTCAGGTGTCCCCTGTCAATATTCTCAAAAG 480
 481 AGAGTCCACCATCTCATAGATGAGATTAGAAAGTGGTCCACAAAAAATATGACACA 540
 541 ACCCATCCATGAACCAATAAAACATGACAGGTCTCATTTCTTCTATTCTTC 600
 601 AAGATAATAATACCTATTAGTGTCTTAAACACCGGCCAACCTTGCAATTCTGTCAATT 660
 661 GGTGACTTTTATTGCCAATTGTGGCTTGAAGGAAATAAAAGGAAAGTCTTTCTTG 720
 721 AACCCATATGGAAGCAATTCAATGAGAGAGATAGAGAGGGATGGAGATTGGGTGG 780
 781 AGAATTGATACGGATCTCTTTAATTGGTATATGTAATCACTCAGAACACGTATACCA 840
 841 TATATGCATCAATGTCATGTCACAGAAAACGTAACTCACGAACACATTCTGTAACATGC 900
 901 ATGCACCAATCATACATTATAACATAGTGTACAGACAATAAAAGATCTTAGTCGAAGA 960
 961 GCATTAGCTGACAGAACAAAAACGTGGATCCCAACCTAAAGAAGGGTATATCTT 1020
 1021 TATTCTATATCTACTTTGATATGACCTAAACCTTGTGTCACCCACAATGTCAGTACG 1080
 1081 ATCGATAATTGTTGACTTGTGTGGATGAGAAATGTATGAGACTGCCATTAGTTTA 1140
 1141 GCCGGATGTGATTGGGTATTGATGACAATATAAGATATATAAAACTGAACAAAACA 1200
 1201 ATTTCTCAACAAATTAAACTACAAGATAATCTCCCTCAGATGATAAAACTAAATGGTAGA 1260
 1261 ATATCGTTGAGTACCCCCAATAATTAAAATCTCCAGCAGAACACTGTGATTCTTCT 1320
 1321 TCGAAGCGAAATTCTCTTCCAAACACCTTAACAAATGTAATTCTGTAGTAAGATT 1380
 1381 AAATTGAAATGATAACACAAGAGTGAATAAAGGTCACTGGTCACCTACTTACCCACTGC 1440
 1441 ACAAAACACACAAGCACACATCCAAAAGTAGTAGTATGATTACACACATTGAAAAAATG 1500
 1501 ACCTCATTATTTAGCCACCTCTCTGTAAAAAAGATTACAAACAAATTACTCCTATCA 1560
 1561 TTATTATAAAATAGTAGCATAACCTCATCTCCAATCCACACCATATAATTACATTATT 1620
 1621 GCCAACATGCTAAAGCTCTTGATTCAGTGAAGGATGTGGTGTCAAATCCAAAGATTC 1680
 1681 TTCATGTGCCCT 1740
 1741 ATCAACTTGAGGGCTTTAGGACCTCTATATAACCTCTCTCAATTGATCATCTCTGATC 1800
 1801 ACACCTCTCAAGCATCTCTCTACTTTCTTGTAACTACACTCCCTTGAGT 1860
 1861 TTCCAATGCCACTGTTGAGGTAAATCAAGTGTATATACATAAAATTATTTGAAAGAT 1920

M A T V E

1921 GATTGATTCAAAGAGAACCCCTTTGTGTTCTTAATAAGATCCATGTATATGAAGTTT 1980
 1981 TAATGTTCATGTTTTTATTGTTAATTGTTAATTAGGCATTGGCAAT 2040
 2041 ATCCCATTTGTGAAAGATCTGTTCTTGGAAAGAGATTAGAATTGTTCTGTCGA 2100
 2101 TTCATCATGAAATCAATCTGGGTCTAGCTTAAATTGTGCTGATCTGACCGACTGTTA 2160
 2161 GATGATTGTTTATATGTAGGCCAATAGAGAGTGTAGTATTCCGAAATAATACAAA 2220
 2221 TCCGAGCAAACATATAATCCTCAATAGTAACCTTGTAATCTCTAAATAATCAAAAAATAAT 2280
 2281 GCTTATTGGGTGATTGGTGTGTTGATGCAGGTTGATCAGCGCAGACAGCATTCCAAG 2340

V V S A Q T A F Q E

2341 AGGAAAAAAAACATGATCAGAAGTAATTACTACAAAAGAGGAAGCTGTAGTAGTAACIG 2400
 E K K H D Q E V I T T K E E A V V V T A
 2401 CACCACCAACATCAGAAACAGCAGAGCCAGCTGAGCTGTTGTCAGGAAAGAGACAA 2460
 P P P S E T A E P A A V V A E E E T T
 2461 CAAAGGAGCAAGAAGAGGCCAGCTATCGGCCAGGAACCTGTGGCCCCAGCTGAAG 2520
 K E Q E E P P A V S A E E P V A P A E V
 2521 TAGAGACAAAGGTGGAAGTTACAGAAGAACCCACAAAGTTGAGGAGAACCGAGAAG 2580
 E T K V E V T E E P P K V E E K P A E V
 2581 TAGAGGAGGCTCCAAAGGAACAGTAGAACAGAACCCAGCTGTTGAGAACCCATCAAGG 2640
 E E A P K E T V E T E P A V E K T I K E

Figure 10 a

2641 AGGAAACTGTAGAGGACTCTGCGTGGCACCTGCTCCGAACCGGAAGCCGAAGTCCCAA 2700
E T V E D S V V A P A P E P E A E V P K
2701 AAGAGAAGGTAATTGCTACTACTGAAACTACTGAGGAAGAAGAAAAAGTGGCAGTTGAAG 2760
E K V I A T T E T E E E E K V A V E E
2761 AAGTTGAAGTGAAGTTGAAACAGAGGGAGGAAGTTACTGAGGAGAAAGACTGAGTAAA 2820
V E V K V E T E G E V T E E K T E *
2821 ATAAGTTGTACAACATTTTATGCACGCCCTTATTTCTCAATTGGAAGTTATAATGTAG 2880
2881 TGGGCTTTGGTAATATTGGGGTTAACATAAGTGGTTAACAGGGTTAACGGCTTTTG 2940
2941 GAATTTAGATATTGGTAAAGGCCTACTTGAACAAAACATAGAAATTGGCACACATGG 3000
3001 GTAAAAGTCAAACTTGTGAGGATGTTCTTGGTTAACATGTGTGCCAAGTAGT 3060
3061 AGAATGTGGTGGTTGTAAAGTTCTCAAGTAGGGTTATGAGTCCTAGTATTATGCT 3120
3121 TGATTGTATGTTGATATGAAAATGGGGTATGTTGGCTTGAAATAAAAGTTTTAATT 3180
3181 ATATAATAAGTGTATTTGTTAACATCATTCTTCATTCTCGGATCAACTACTGAT 3240
3241 CATGCCCTGGTAAGCTATTGCCCTACCAACTAGCTAACATCGAACGCGAGCCC 3292

Figure 10b